

Organization Applicant

 Street : 351 Tache Avenue
 City : Winnipeg
 State : Manitoba
 Country : Canada
 PostalCode : R2H 2A6
 PhoneNumber : 204-235-3206
 FaxNumber :
 EmailAddress :

<110> OrganizationName : St. Boniface General Hospital Research Centre

Individual Applicant

 Street : 351 Tache Avenue
 City : Winnipeg
 State : Manitoba
 Country : Canada
 PostalCode : R2H 2A6
 PhoneNumber :
 FaxNumber :
 EmailAddress :

<110> LastName : Mesaeli
 <110> FirstName : Nasrin
 <110> MiddleInitial :
 <110> Suffix :

Application Project

 <120> Title : Transgenic Mouse over-expressing calreticulin (CRT) in vascular
 smooth muscle cells
 <130> AppFileReference : 81190-2602
 <140> CurrentAppNumber :
 <141> CurrentFilingDate : ____-____-____

Earlier Applications

 <150> PriorAppNumber : US 60/455399
 <151> PriorFilingDate : 2003-03-18

Sequence

 <213> OrganismName : Artificial
 <400> PreSequenceString :
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 gaaggggcca ggggtgactt gctgctaaac aaggcactcc ctagagaagc acccgctaga 660
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crt sequences.workFile

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<212> Type : DNA

<211> Length : 2655

SequenceName : SM22a-CRT

SequenceDescription :

Feature

Sequence: SM22a-CRT:

<221> FeatureKey : CDS

<222> LocationFrom : 1319

<222> LocationTo : 2653

Other Information : CRT coding sequence

CDSJoin : No

Sequence

<213> OrganismName : Artificial

<400> PreSequenceString :

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                                crt sequences.workFile
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```

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<212> Type : DNA
<211> Length : 2691
      SequenceName : sm22a-crt-ha
      SequenceDescription :

```

Feature

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Sequence: sm22a-crt-ha:
<221> FeatureKey : CDS
<222> LocationFrom : 1319
<222> LocationTo : 2689
      Other Information :
      CDSJoin : No

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Feature

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Sequence: sm22a-crt-ha:
<221> FeatureKey : misc_feature
<222> LocationFrom : 2635
<222> LocationTo : 2670
      Other Information : HA insertion
      CDSJoin : No

```

Sequence

```

-----
<213> OrganismName : Artificial
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DMHGDSEYNI MFGPDICGPG TTKVHVIFNY KGKNVLINKD IRCKDDEFTH LYTLIVRPDN 180
TYEVKIDNSQ VESGSLEDDW DFLPPKKIKD PDASKPEDWD ERAKIDDPTD SKPEDWDKPE 240
HIPDPDAKKP EDWDEEMDGE WEPPVIQNPE YKGEWKPRQI DNPDKGTWI HPEIDNPEYS 300
PDANIYAYDS FAVLGLDLWQ VKSGTIFDNF LITNDEAYAE EFGNETWGV TAEKQMKDK 360
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```
<212> Type : PRT
```

crt sequences.WorkFile

<211> Length : 418
SequenceName : crt
SequenceDescription :

Sequence

<213> OrganismName : Artificial
<400> PreSequenceString :
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EKDKGLQTSQ DARFYALSAR FEPFSNKGQP LVVQFTVKHE QNIDCGGGYV KLFPAGLDQK 120
DMHGDSEYNI MFGPDICGPG TKKVHVIFNY KGKNVLINKD IRCKDDEFTH LYTLIVRPDN 180
TYEVKIDNSQ VESGSLEDDW DFLPPKKIKD PDASKPEDWD ERAKIDDPDPTD SKPEDWDKPE 240
HIPDPDAKKP EDWDEEMDGE WEPPVIQNP EYKGEWKPRQI DNPDYKGTWI HPEIDNPEYS 300
PDANIYAYDS FAVLGLDLWQ VKSGTIFDNF LITNDEAYAE EFGNETWGV T KTAEQMKDK 360
QDEEQRLKEE EEEKKRKEE EAEDEEDKD DKEDEDEE DKDEEEEEEAA AGLEYPYDVP 420
DYARQAKDEL 430
<212> Type : PRT
<211> Length : 430
SequenceName : crtha
SequenceDescription :

Feature

Sequence: crtha:
<221> FeatureKey : MISC_FEATURE
<222> LocationFrom : 413
<222> LocationTo : 424
Other Information : HA Tag
CDSJoin : No